



Wave Life Sciences

Corporate Presentation

April 28, 2026

Forward-looking statements

This document contains forward-looking statements. All statements other than statements of historical facts contained in this document, including statements regarding possible or assumed future results of operations, preclinical and clinical studies, business strategies, research and development plans, collaborations and partnerships, regulatory activities and timing thereof, competitive position, potential growth opportunities, use of proceeds and the effects of competition are forward-looking statements. These statements involve known and unknown risks, uncertainties and other important factors that may cause the actual results, performance or achievements of Wave Life Sciences Ltd. (the “Company”) to be materially different from any future results, performance or achievements expressed or implied by the forward-looking statements. In some cases, you can identify forward-looking statements by terms such as “may,” “will,” “should,” “expect,” “plan,” “aim,” “anticipate,” “could,” “intend,” “target,” “project,” “contemplate,” “believe,” “estimate,” “predict,” “potential” or “continue” or the negative of these terms or other similar expressions. The forward-looking statements in this presentation are only predictions. The Company has based these forward-looking statements largely on its current expectations and projections about future events and financial trends that it believes may affect the Company’s business, financial condition and results of operations. These forward-looking statements speak only as of the date of this presentation and are subject to a number of risks, uncertainties and assumptions, including those listed under Risk Factors in the Company’s Form 10-K and other filings with the SEC, some of which cannot be predicted or quantified and some of which are beyond the Company’s control. The events and circumstances reflected in the Company’s forward-looking statements may not be achieved or occur, and actual results could differ materially from those projected in the forward-looking statements. Moreover, the Company operates in a dynamic industry and economy. New risk factors and uncertainties may emerge from time to time, and it is not possible for management to predict all risk factors and uncertainties that the Company may face. Except as required by applicable law, the Company does not plan to publicly update or revise any forward-looking statements contained herein, whether as a result of any new information, future events, changed circumstances or otherwise.



Our Mission

To unlock the broad
potential of RNA medicines
to transform human health



Building a leading RNA medicines company

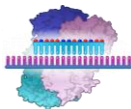
Differentiated RNA medicines platform and chemistry



- **Proprietary chemistry**
- Leveraging deep insights in **human genetics**
- Strong and **broad IP**
- **In-house GMP** manufacturing

Translating genetic insights into potentially best-in-class medicines

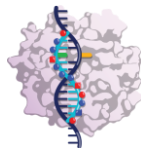
RNAi



WVE-007 (obesity)

- Differentiated mechanism focused on fat loss and muscle preservation

RNA editing



WVE-006 (AATD)

WVE-008 (liver disease)

- Restoration of functional protein production

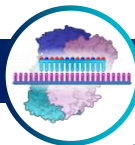
Other modalities: **Splicing, antisense silencing**

Unlocking emerging pipeline

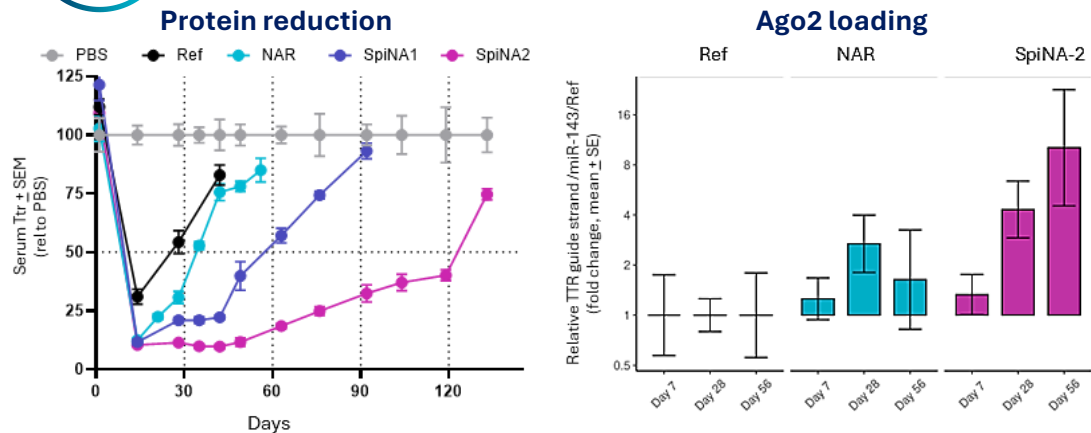
- **Extra-hepatic capabilities:** with RNAi and RNA editing
- **Bifunctional modalities:** single oligonucleotide constructs for dual RNAi silencing or RNAi silencing + RNA editing

Well capitalized with expected cash runway into 3Q 2028

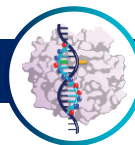
For over a decade Wave has been extending the frontiers of RNA therapies delivering breakthroughs in nucleic acid chemistry



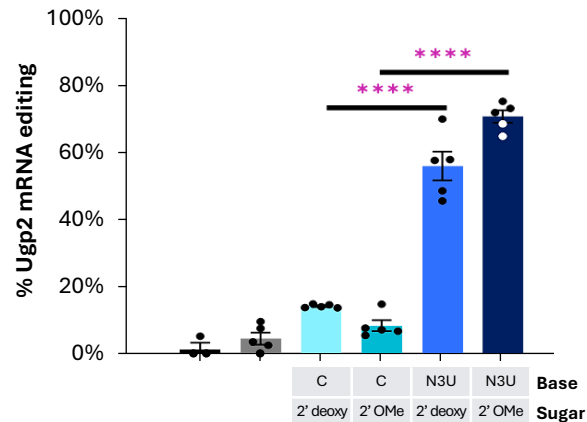
RNAi — SpiNA



Substantial improvements in potency, duration of activity, and Ago2 loading with Wave's proprietary SpiNA design



RNA editing — AIMer



Increased RNA editing efficiency achieved with proprietary chemistry

Proprietary chemistry has dramatically increased potency and durability

Robust RNA medicines pipeline with first-in-class RNAi and RNA editing programs

Program	Discovery	IND / CTA Enabling Studies	Clinical	Patient population (US & Europe)
RNAi (SpiNA)				
WVE-007 (GalNAc) INHBE (obesity)	<div></div>			175M (>1 billion globally)
GalNAc / extra-hepatic Multiple	<div></div>			--
RNA EDITING (AIMer)				
WVE-006 (GalNAc) SERPINA1 (AATD)	<div></div>			200K
WVE-008 (GalNAc) PNPLA3 (liver disease)	<div></div>			9M
GalNAc / extra-hepatic Multiple	<div></div>			--
SPLICING				
WVE-N531 Exon 53 (DMD)	<div></div>			2.3K
Other exons (DMD)	<div></div>			Up to 18K
ALLELE-SELECTIVE SILENCING				
WVE-003 mHTT (HD)	<div></div>			25K Symptomatic (SNP3) 60K Pre-Symptomatic (SNP3)

WVE-007
GalNAc-siRNA silencing

Obesity

WVE-007 (investigational INHBE GalNAc-siRNA) is a potentially transformative approach for the > 1 billion people living with obesity globally

Significant unmet need in obesity

Current standard of care: Focused on caloric restriction by reducing appetite and slowing gastric emptying

Incretins limited by:

- ✗ Muscle loss¹
- ✗ Frequent dosing²
- ✗ Poor tolerability³
- ✗ High discontinuation rates^{4,5}



Resulting in need for therapies that can:

- Induce **fat loss** with **muscle preservation**
- Leverage **orthogonal mechanism** for **enhanced** efficacy and **maintain metabolic improvements** after incretin cessation

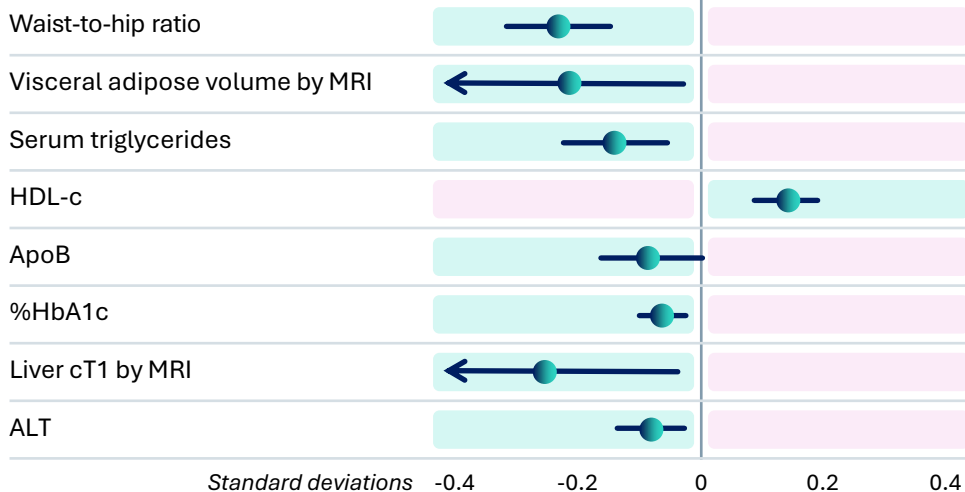
WVE-007

Focused on adipocyte lipolysis and not caloric deficit

- ✓ Drives total and visceral fat loss
- ✓ Preserves muscle
- ✓ Potential 1–2x per year dosing
- ✓ Generally safe and well tolerated

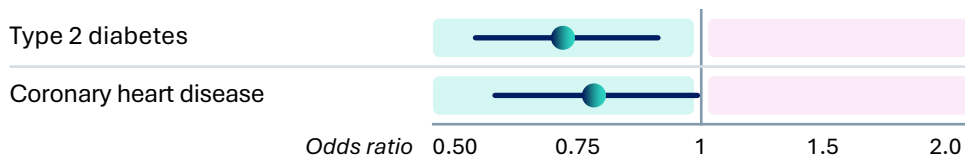
Human genetic data demonstrate that heterozygous INHBE loss-of-function (LoF) carriers have lower visceral fat and a healthier metabolic profile

Heterozygous INHBE LoF carriers versus non-carriers:

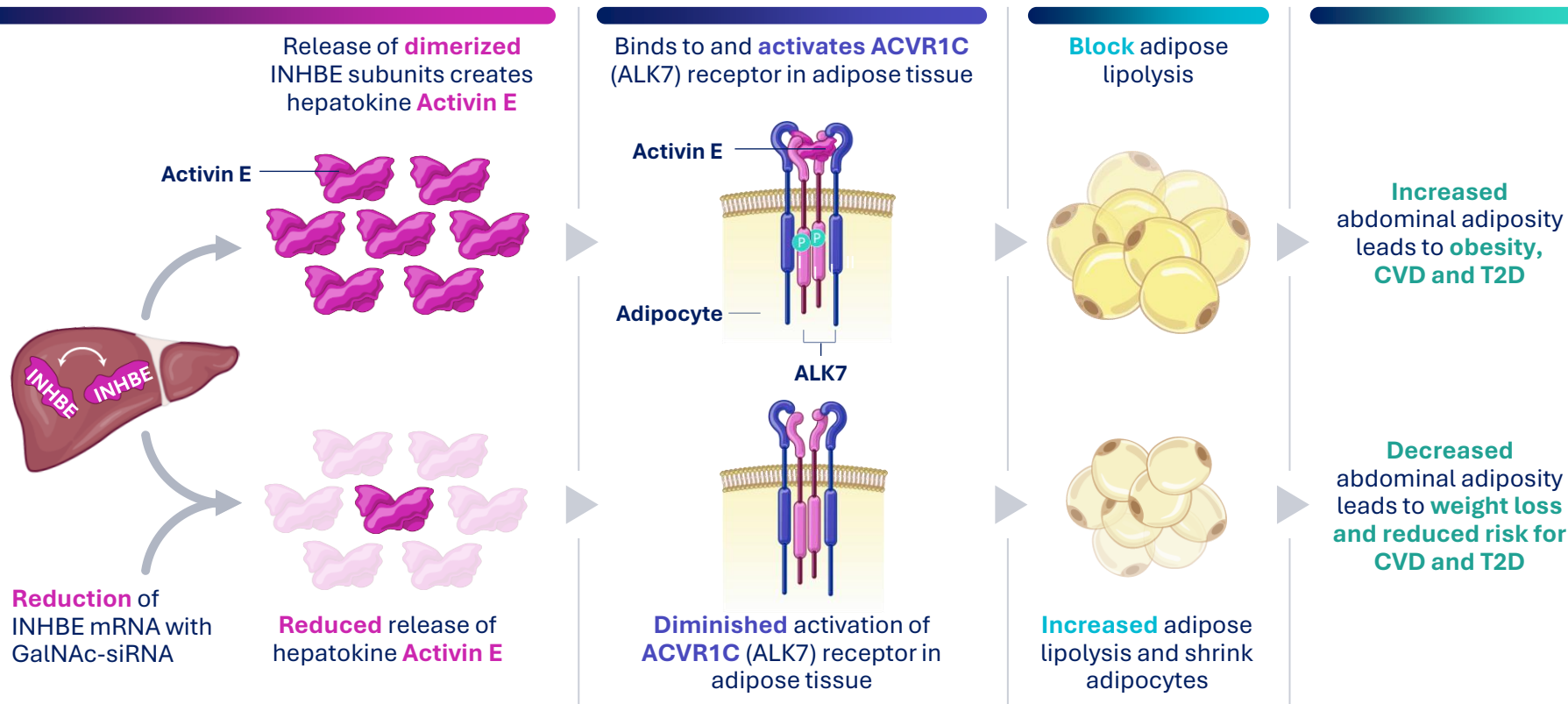


Favorable traits :

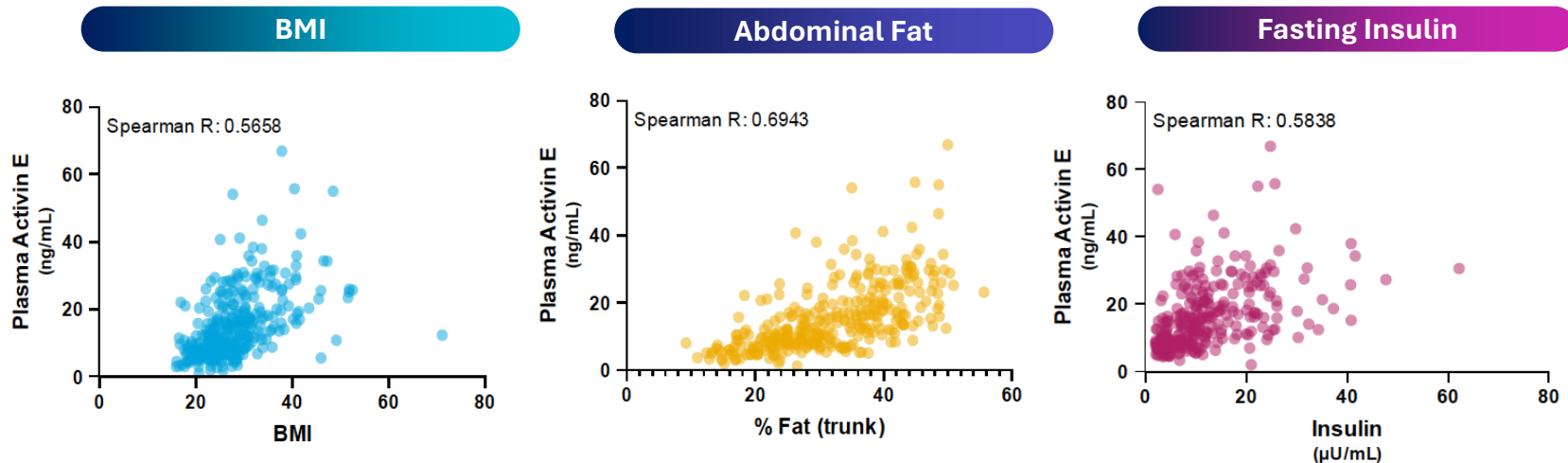
- ✓ Lower abdominal obesity
- ✓ Reduced visceral fat
- ✓ Lower triglycerides
- ✓ Higher “good” cholesterol
- ✓ Lower ApoB
- ✓ Improved glucose control
- ✓ Less liver inflammation/fibrosis
- ✓ Less liver damage



Silencing INHBE mRNA has the potential to treat obesity and associated metabolic diseases

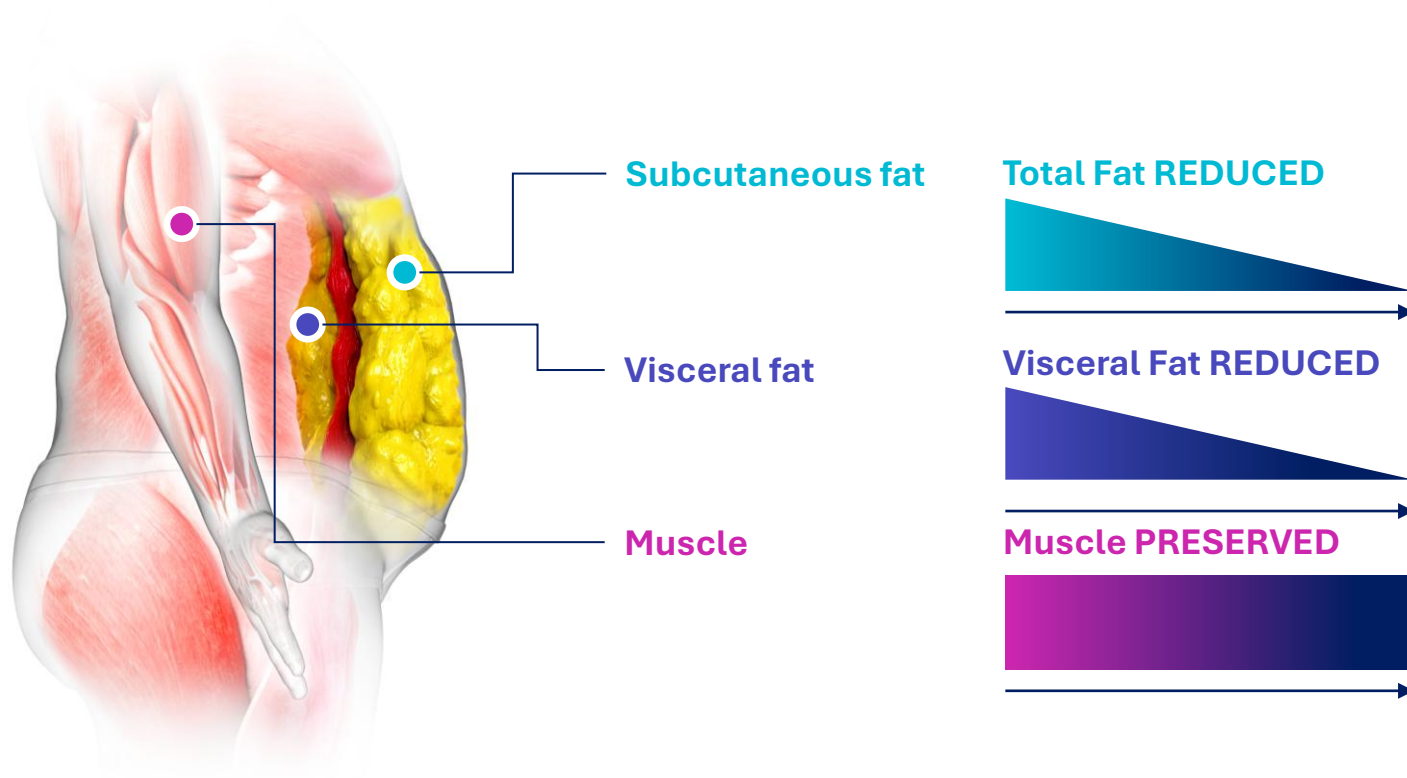


Higher circulating Activin E levels are correlated with higher BMI, higher abdominal fat, and higher fasting insulin in non-diabetic adults



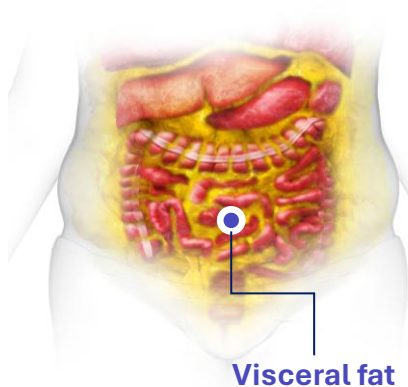
Further supports INHBE suppression as a weight loss approach for individuals living with obesity

WVE-007 is designed to improve body composition by decreasing fat and preserving muscle



WVE-007's mechanism is focused on lipolysis and directly reducing visceral and subcutaneous fat

Visceral fat drives insulin resistance and many cardiometabolic disorders



Waist circumference is a clinical proxy for visceral fat

- MASH
- Type 2 diabetes
- Cardiovascular diseases
- PCOS

Reduced visceral fat is associated with multiple health benefits

Visceral fat decrease

≥ 5%

Improved insulin sensitivity¹: Lower HbA1c and better lipid profile

≥ 5-10%

Reduced cardiovascular risk²: Reduced blood pressure, improved lipids, lower systemic inflammation

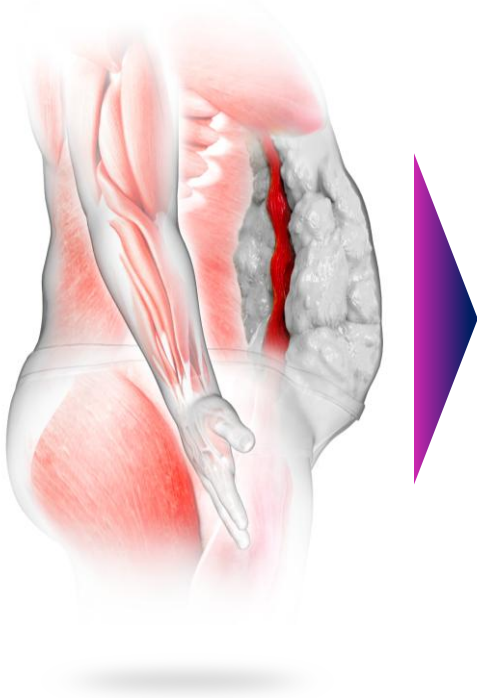
≥ 10%

Reduced liver fat (steatosis)³: Significant reduction in hepatic triglycerides, improved liver enzymes

Decreased hepatic fibrosis⁴: Resolution of steatohepatitis in up to 90%, fibrosis regression in many cases

WVE-007 aims to address a key limitation of current standard of care: up to 40% of weight loss is driven by muscle loss

Preservation of muscle mass is linked to many health benefits

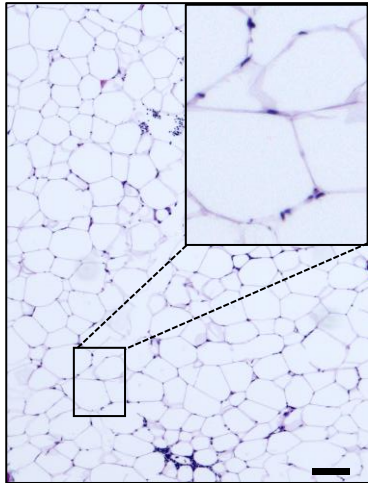


- Higher basal metabolic rate (BMR)¹
- Improved insulin sensitivity^{2,3}
- Increased caloric expenditure post-exercise¹
- Preserve muscle strength and function⁴
- Reduced visceral fat^{5,6}
- Prevent weight regain^{7,8}
- Improved glucose homeostasis^{2,3}
- Increased bone density, strength, function, and longevity^{9,10}

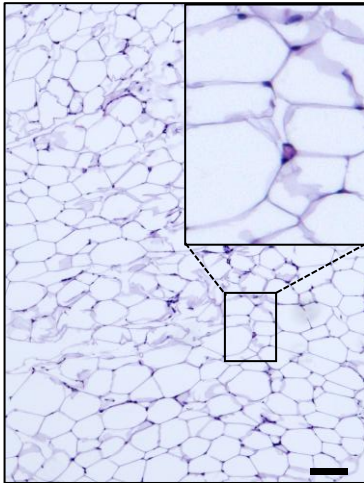
A single dose of INHBE GalNAc-siRNA led to shrinkage of adipocytes in DIO mice

DIO

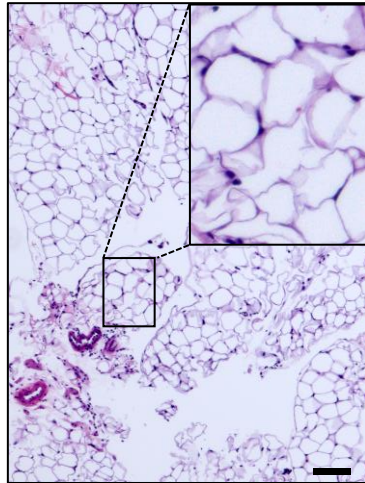
PBS



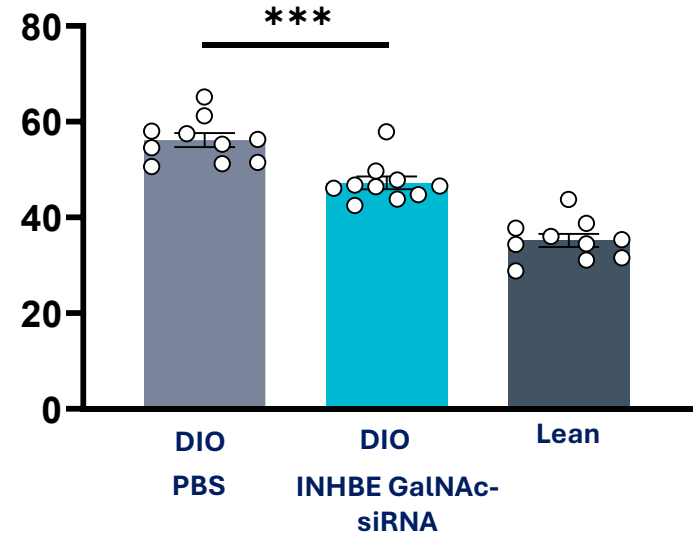
INHBE GalNAc-siRNA



Lean

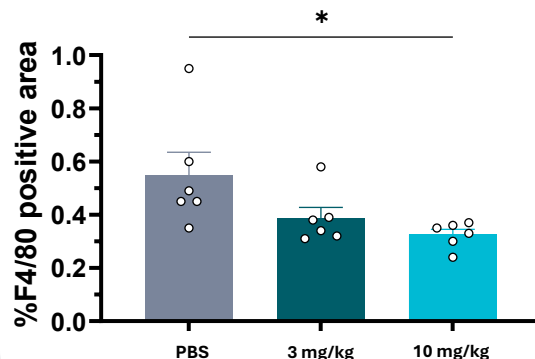
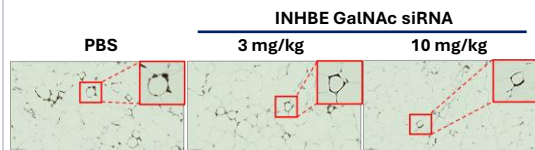


**Mean adipocyte diameter
(μm)**

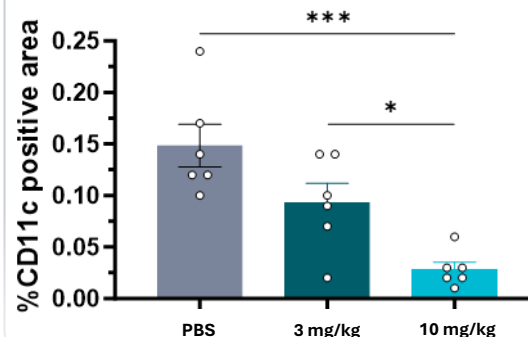
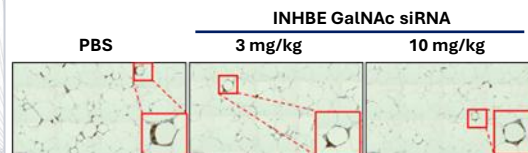


A single dose of INHBE siRNA led to a lower inflammatory state of visceral adipose tissues in DIO mice, with strong suppression of pro-inflammatory M1 macrophages in visceral fat

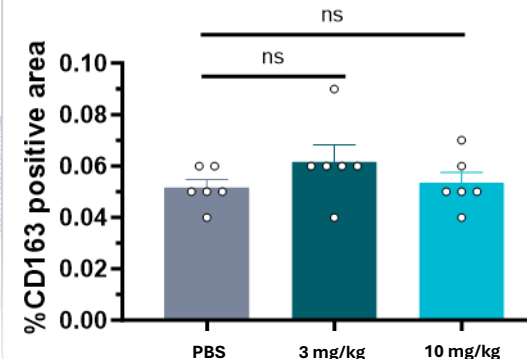
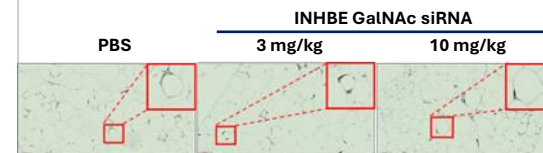
Macrophages (M ϕ) (F4/80)



Pro-inflammatory (M1) M ϕ (CD11c)

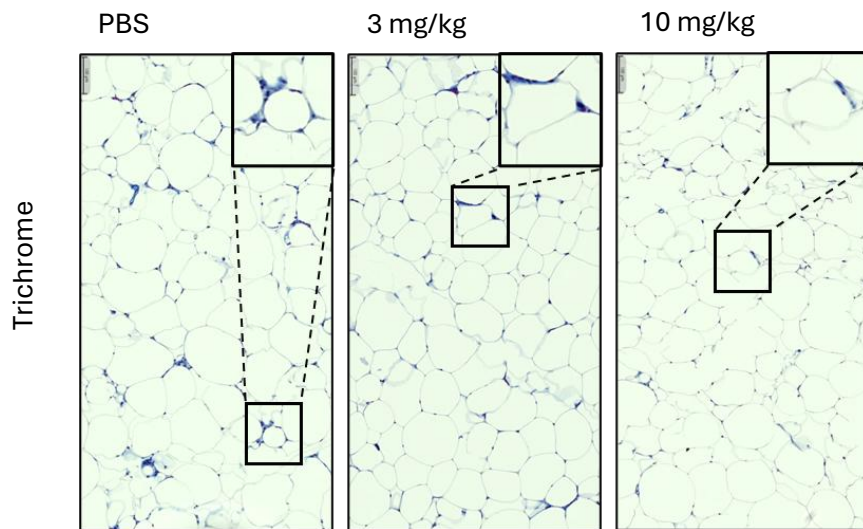


Anti-inflammatory (M2) M ϕ (CD163)

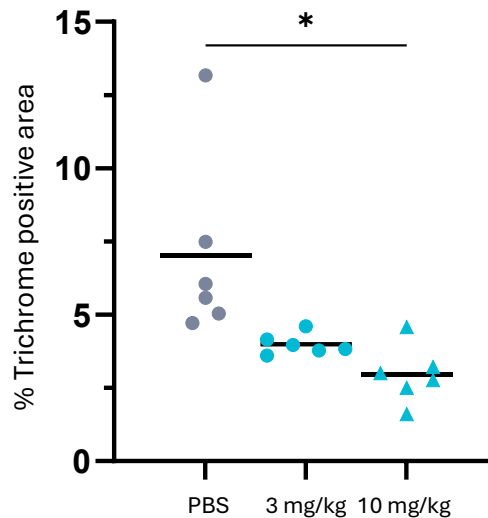


Lowering of inflammatory state of epiWAT visceral fat induced by single dose of INHBE siRNA resulted in 58% reduction of adipose fibrosis

Reduced staining illustrates decreased tissue fibrosis



Fibrosis in mouse adipose (Day 56)




Treatment with WVE-007 (investigational INHBE GalNAc-siRNA) is expected to drive fat reduction and improve key measures of cardiometabolic health

Leading siRNA design (SpiNA)¹

Driving fat reduction

And improving clinical outcomes

WVE-007 (INHBE GalNAc-siRNA)

✓  Proprietary, clinically validated chemistry

✓ Subcutaneous delivery (GalNAc)

✓ Potential for infrequent dosing (1 – 2x year)

↓ Reduction of INHBE mRNA and circulating Activin E



Adipocyte lipolysis



Adipocyte size



Proinflammatory macrophages



Fibrosis



Insulin sensitivity

Cardiometabolic outcomes

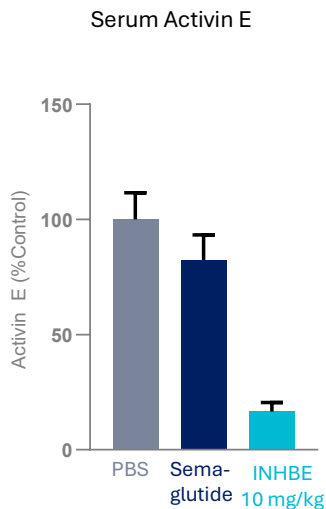
↓ Risk of CVD

↓ Risk of T2D

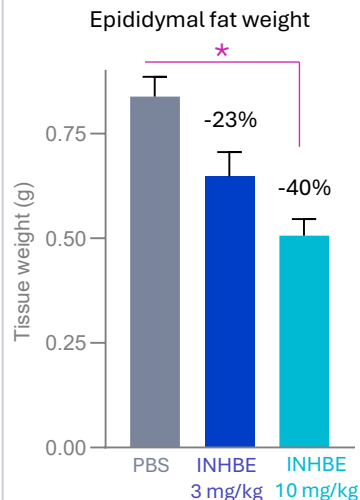
Single dose of INHBE GalNAc-siRNA led to durable Activin E reductions, and sustained improvements in body composition in DIO mice



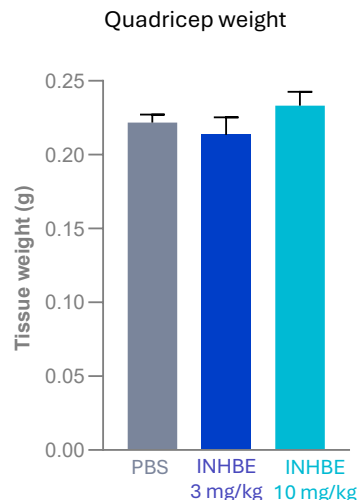
Durable Activin E reduction



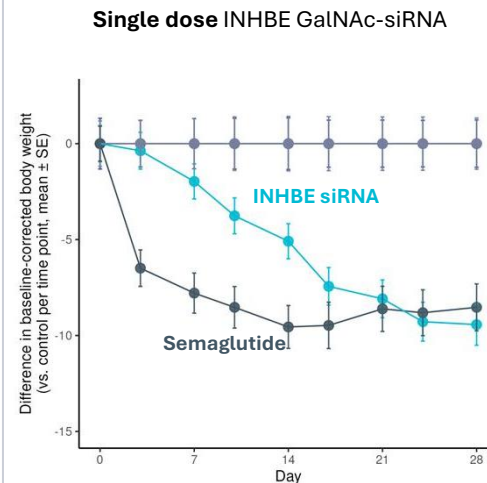
Reduction in fat



Muscle preservation



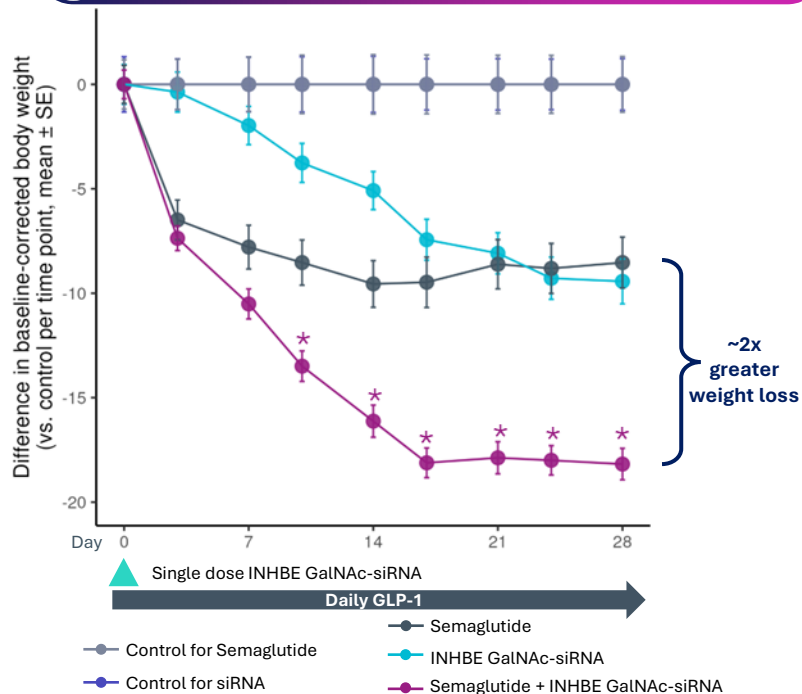
Reduction in body weight



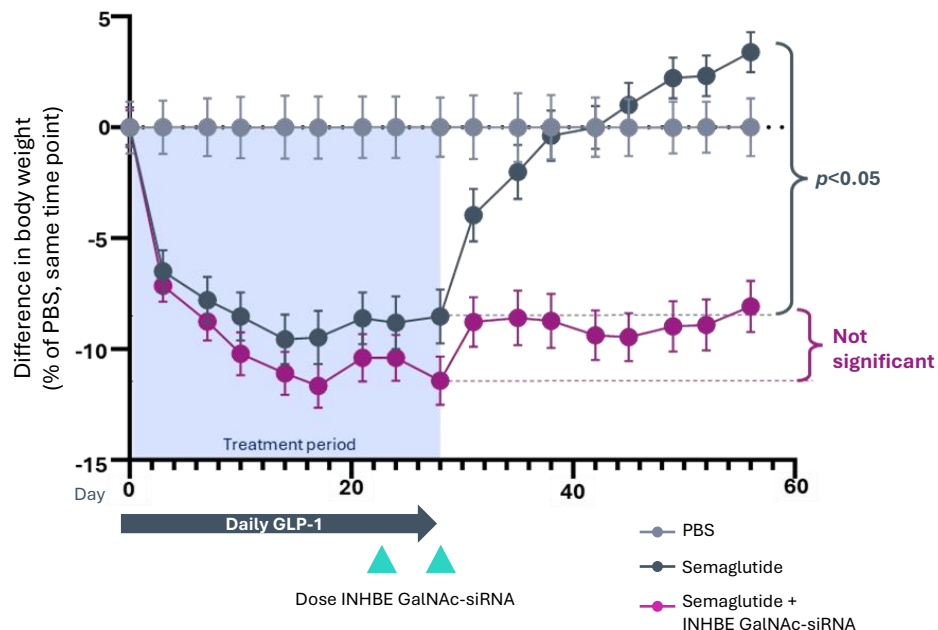
WVE-007 has potential for use synergistically with GLP-1s or to curtail weight regain after the cessation of treatment with GLP-1, based on preclinical data



Combined with GLP-1: Greater weight loss



After cessation of GLP-1: Curtails weight regain



Phase 1 portion of INLIGHT trial is investigating WVE-007 in individuals living with overweight or obesity, otherwise healthy



Phase 1 lower BMI (SAD)

- **Average BMI: 32 kg/m²**
- **HbA1c: <5.9%**
- **Otherwise healthy**

- Assessments include: Safety and tolerability, PK, Activin E, body composition (DEXA), biomarkers, body weight
- No diet or exercise modifications

SAD Cohort 4
600 mg (n=32)

SAD Cohort 3
400 mg (n=32)

SAD Cohort 2
240 mg (n=32)

SAD Cohort 1
75 mg (n=8)
PK/PD and safety only (no DEXA)

Evaluate safety, tolerability, and PK

Phase 2a high BMI (MAD)

MAD high BMI (35–50 kg/m²)+T2D cohorts

MAD high BMI (35–50 kg/m²) cohorts

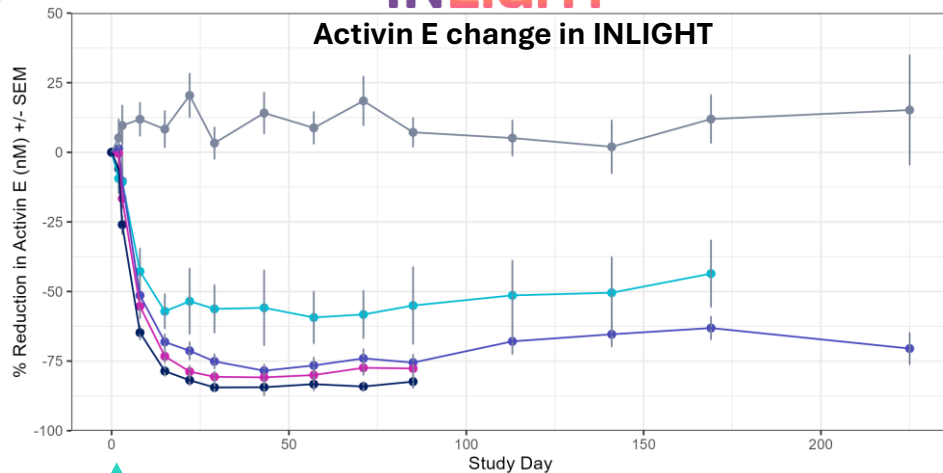
- Individuals with **higher BMI with and without type 2 diabetes**
- Assessments to additionally include:
 - Body composition (MRI in addition to DEXA)
 - Liver fat (MRI-PDFF)
- Diet and exercise modifications included

**Evaluate safety, tolerability, and PK;
assess metabolic and body composition
improvements as well as weight loss**

Clinically meaningful improvements in body composition at six months following a single dose of WVE-007



Activin E change in INLIGHT



Phase 1 otherwise healthy (SAD) Lower BMI of ~32 kg/m²

Improved body composition six months following single 240 mg dose:

- Significant reduction in **visceral fat** (-14%*)
 - Reduction in **waist circumference** (-3%)
 - Reduction in **total fat** (-5%)
 - Stabilization of **lean mass** (+2%)
 - Reduction in **body weight** (-1%)
- 400 mg three-month data emphasize impact of baseline body composition on therapeutic effect
 - **Durable and dose-dependent** suppression of Activin E sustained through at least 7 months continues to support **1-2x yearly dosing**
 - **Generally safe and well tolerated**

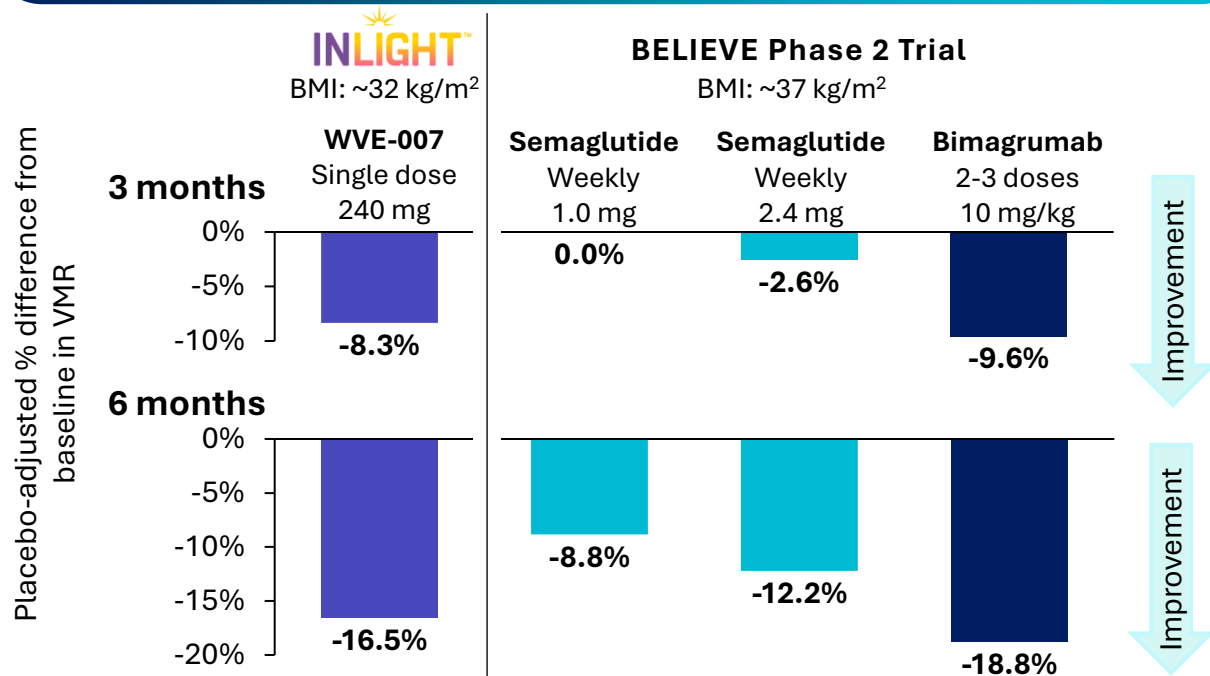
Durability of suppression continues to support dosing WVE-007 once or twice per year

Single dose of WVE-007 in a lower BMI population led to greater improvement in body composition by VMR versus semaglutide

Visceral Fat-to-Muscle Ratio (VMR)

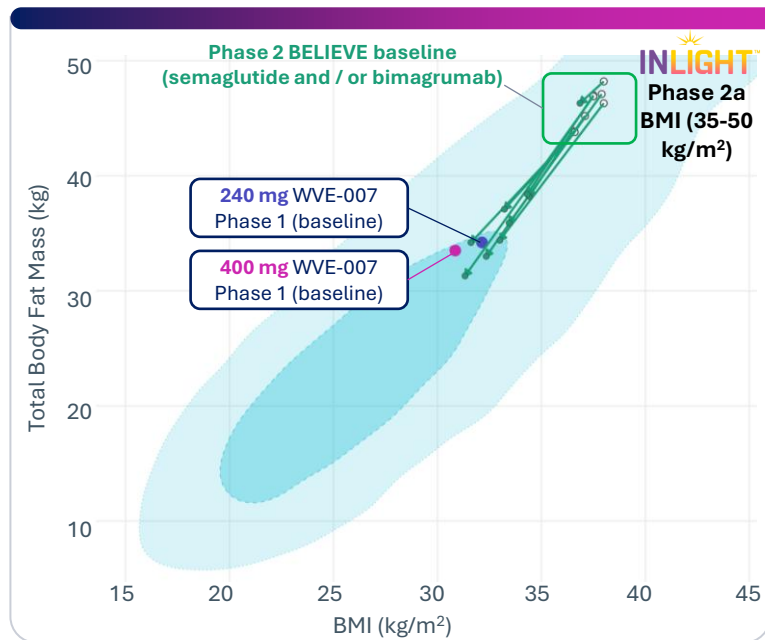
- Established measure of body composition integrating harmful visceral fat and beneficial lean mass in a single index
- Lower VMR is associated with decreased risk of MASH / MAFLD,^{1,2} type 2 diabetes,³ and cardiometabolic disorders (e.g., dyslipidemia, hypertension)^{1,3}

Improvement in body composition by VMR at 3 months and 6 months⁴

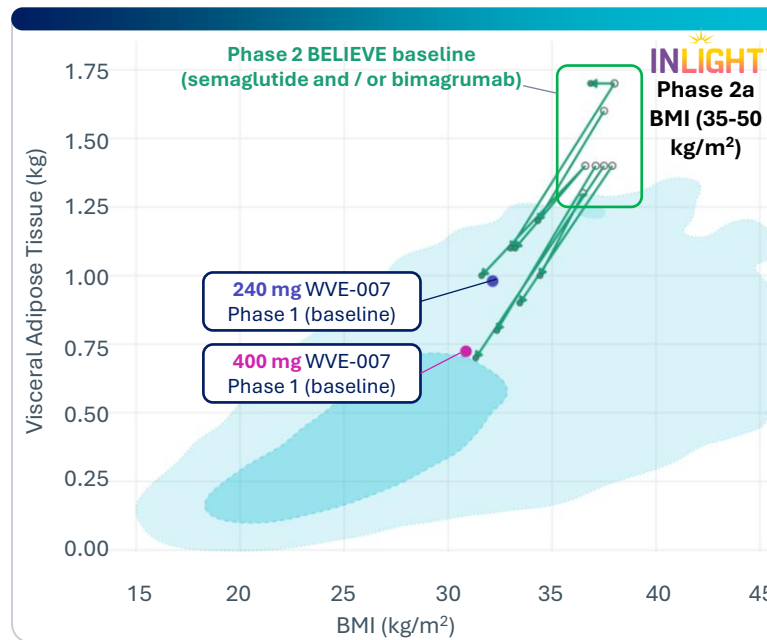


INLIGHT Phase 2a: higher BMI population aligns with Phase 2 and 3 obesity trials, potential for continued improvements in body composition

Total fat and BMI at baseline



Visceral fat and BMI at baseline



BELIEVE Phase 2 study

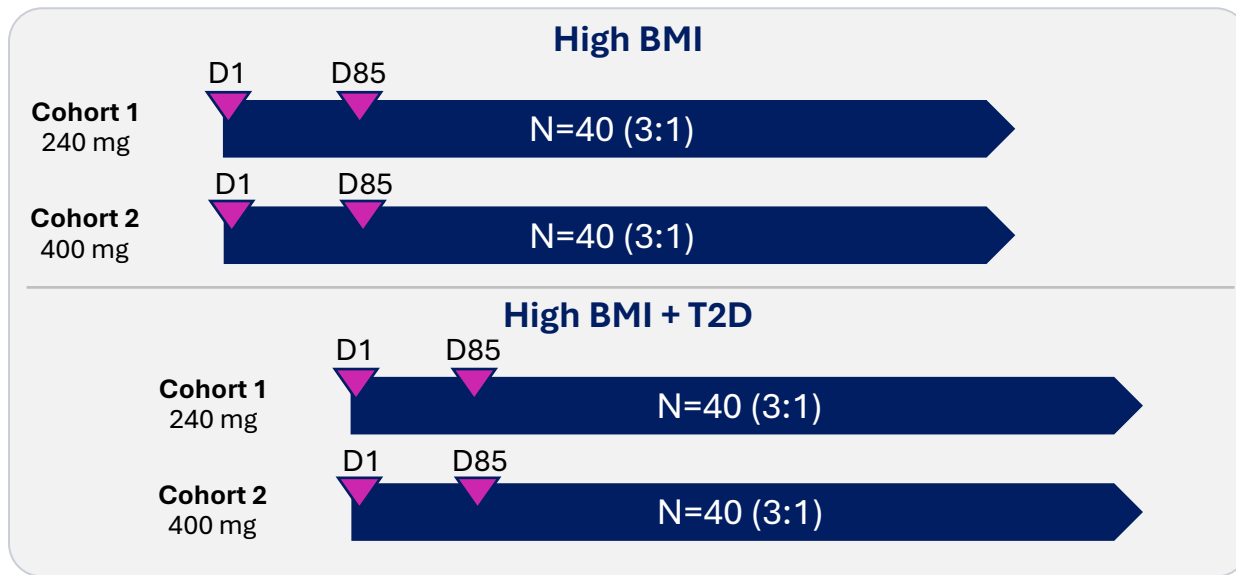
- Baseline (by cohort)
- Week 48 (by cohort)

US population data from NHANES (VAT and BMI)

- 50% of population
- 95% of population

Expect greater fat loss in INLIGHT Phase 2a in participants with more excess fat

High BMI Phase 2a (MAD) global, placebo-controlled study will inform further development in obesity, as well as MASH, type 2 diabetes, and CVD



- Individuals with **higher BMI** (35–50 kg/m²) and **comorbidities**
- Assessments include:
 - Body weight
 - Body composition (MRI in addition to DEXA)
 - Liver fat (MRI-PDFF)
 - HbA1c, lipid levels, CRP
 - Muscle function
- Diet and exercise modifications included
- 12-month study duration
 - First assessment Day 85 (3 months)

Expect to initiate high BMI Phase 2a (MAD) portion of INLIGHT in 2Q 2026

On track to initiate the Phase 2a portion of INLIGHT in 2Q 2026; combination and maintenance studies of WVE-007 expected to initiate in 2026

Monotherapy

Single agent in individuals living with obesity



- To induce fat loss with muscle preservation and favorable safety and tolerability

Combination

Add-on to incretin treatments

- To leverage an orthogonal mechanism to incretins for enhanced efficacy

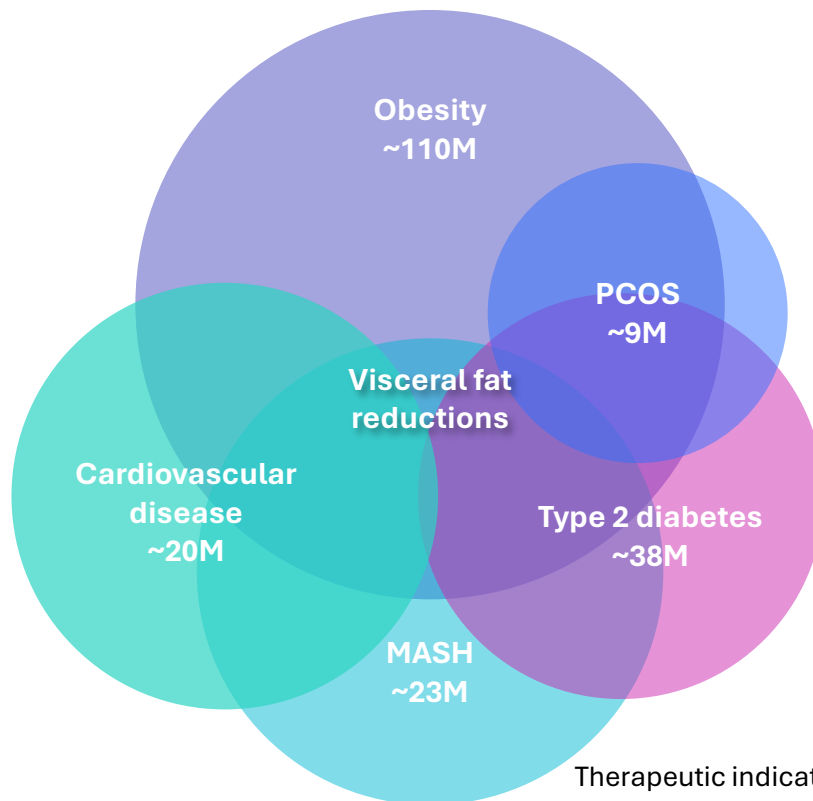
Maintenance

An off-ramp post-incretin treatments

- To prevent weight regain and maintain metabolic improvements upon incretin cessation

Potential to address more than one billion individuals with obesity globally

INHBE silencing provides opportunity to address additional significant indications through lowering of visceral fat



Therapeutic indications and US patient populations

WVE-006

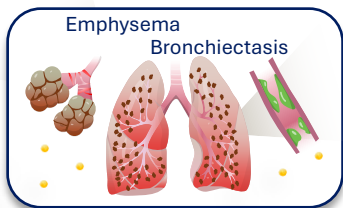
RNA editing (AlMer)

Alpha-1 antitrypsin deficiency (AATD)

AATD impacts multiple organ systems and has limited treatment options

- AATD is a rare, inherited genetic disorder that is commonly caused by a G-to-A point mutation in the SERPINA1 gene
- Pi*ZZ genotype is leading cause of severe AATD, predisposing to progressive lung damage, liver damage or both
- Aggregation of mutant Z-AAT protein in hepatocytes and a lack of functional, wild-type M-AAT drives liver and lung disease, respectively

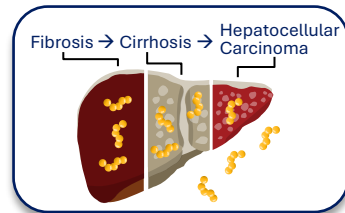
AATD Lung Disease



- **Treatment goal:** Minimize episodic exacerbations and associated damage
- Lung damage occurs during exacerbations that induce an inflammatory acute phase response, when more AAT protein is needed for protection

- **Weekly IV augmentation therapy is only treatment option**
 - No protective increase in AAT protein levels during acute phase response without additional IV infusions

AATD Liver Disease



- **Treatment goal:** Decrease Z-AAT protein
- Progressive liver disease results from Z-AAT-induced proteotoxic stress

- **No approved therapies**

~200K people in the US and Europe are homozygous for the Z allele (Pi*ZZ genotype)

WVE-006: Potential first-in-class, convenient therapy for AATD that addresses both liver and lung manifestations of the disease



WVE-006 (RNA editing)



Proprietary chemistry



Highly specific (no bystanders)



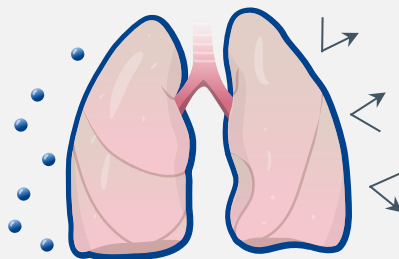
Subcutaneous delivery (GalNac)



Infrequent dosing



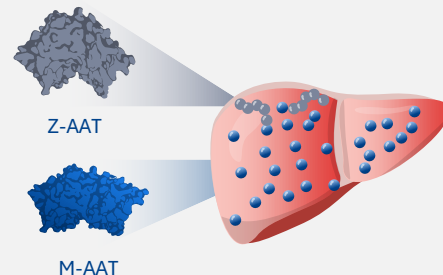
Restore circulating M-AAT and physiological AAT protein production



M-AAT reaches lungs to protect from proteases and **reduce risk of lung pathology**



Reduce Z-AAT protein aggregation in liver



RNA correction replaces mutant Z-AAT protein with wild-type M-AAT protein to **reduce risk of liver pathology**

RNA editing aims to increase M-AAT and restore physiological AAT production during acute phase response

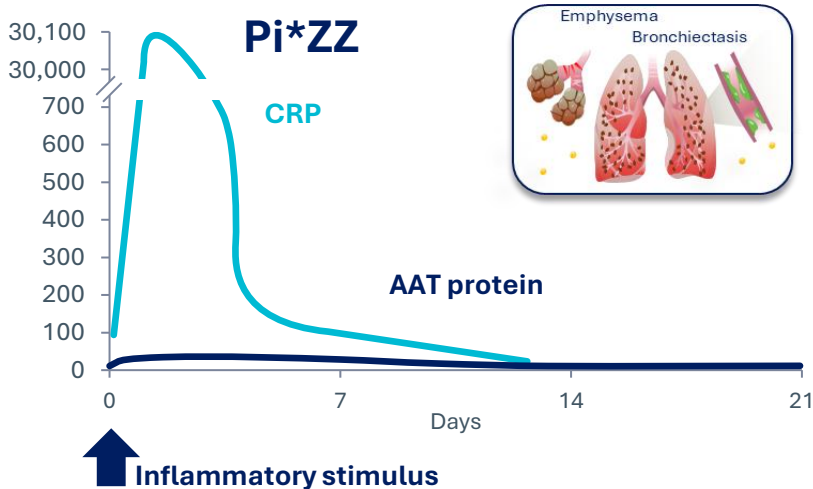
Genotype	Null	Pi*ZZ	Pi*MZ	Pi*MM
	No AAT protein	100% Z-AAT	Z-AAT and M-AAT	100% M-AAT
AAT levels increase during acute phase response	No	No	Yes	Yes
Risk of lung disease	Very high	High	Low	Normal
Risk of liver disease	None	High	Low	Normal

>50% RNA editing
>11 μ M AAT

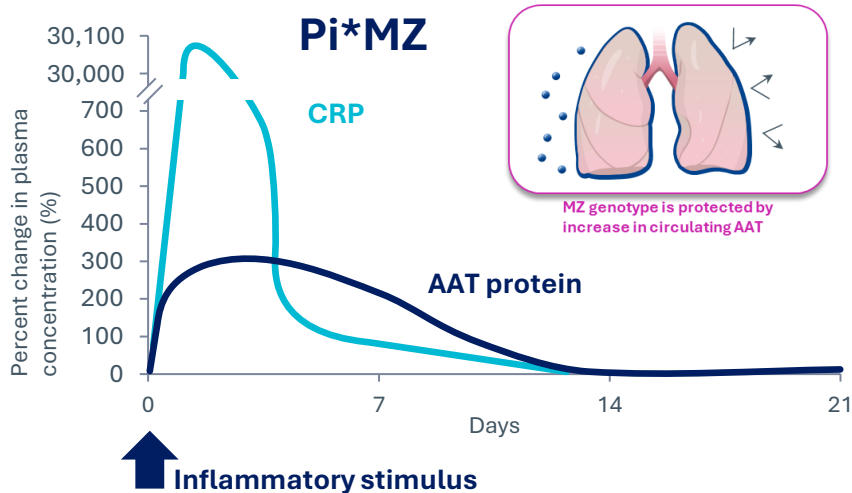
Goal: Shift Pi*ZZ individuals to AAT biomarker profile consistent with Pi*MZ genotype

RNA editing aims to restore production of dynamic and therapeutically relevant levels of AAT protein in Pi*ZZ individuals during acute phase response

Lung damage occurs during exacerbations, when more AAT protein is needed for protection



AAT protein has protective functions and is produced during acute phase response

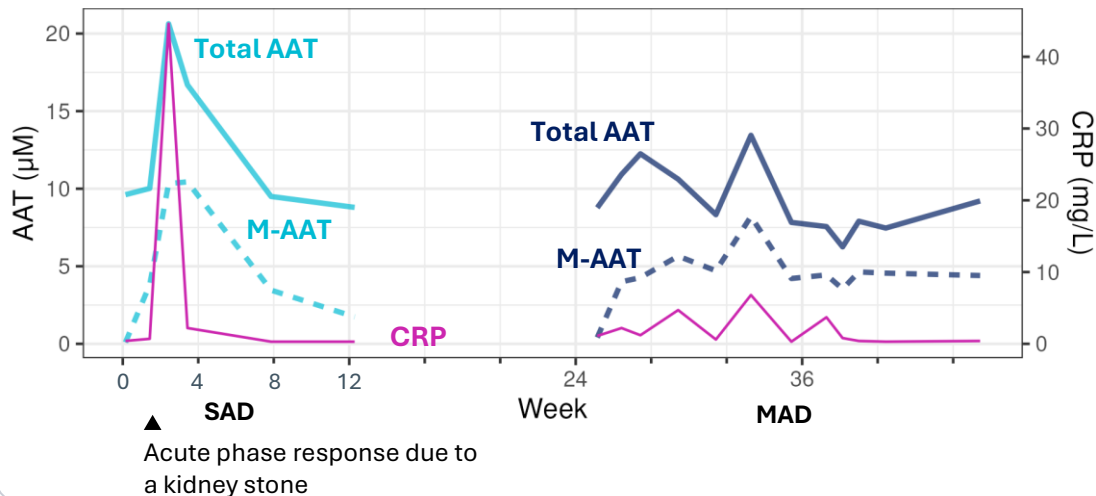


RNA editing has potential to restore dynamic AAT response to inflammation

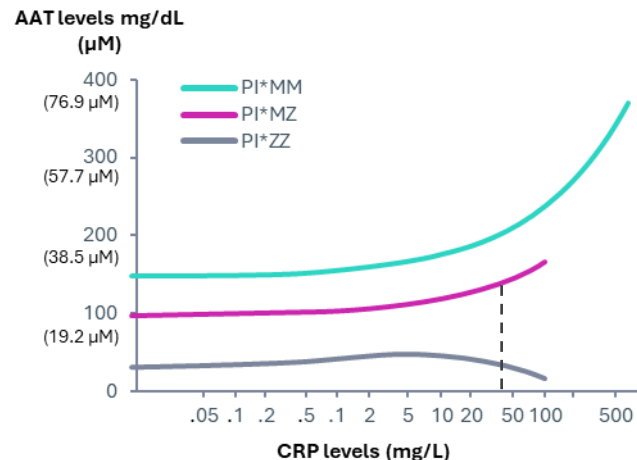
First-ever demonstration of ability to restore physiological serum AAT production; total AAT reached 20.6 μM during acute phase response

Pi*ZZ patients have a reduced capacity to produce AAT protein during an acute phase response

Following WVE-006 200 mg single dose, total AAT and M-AAT increased significantly in one patient during an acute phase response



Published data¹ on CRP levels and AAT levels across different genotypes

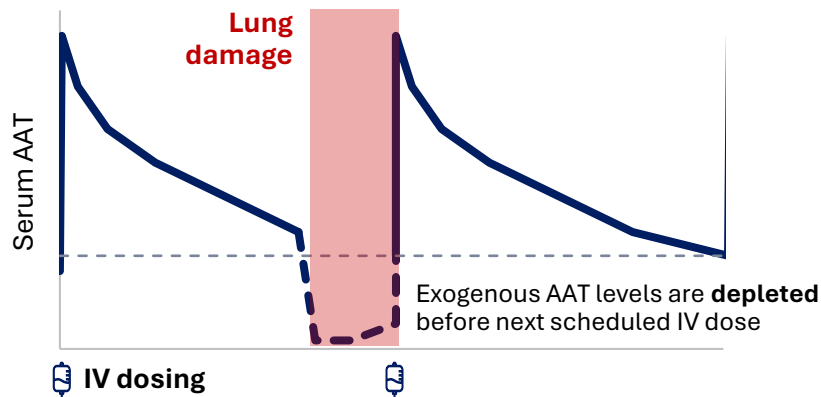


AAT response in Pi*ZZ participant treated with WVE-006 mirrors Pi*MZ phenotype

WVE-006 enables endogenous AAT production during an acute phase response while augmentation therapy may leave patients at risk

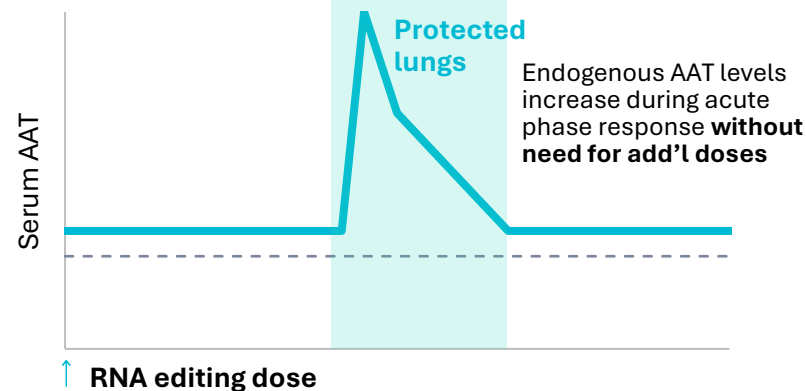
Illustrative model of impact of acute phase response

Augmentation therapy



- Augmentation therapy has **no impact** on liver disease

WVE-006 treatment approach



- WVE-006 also reduces levels of Z-AAT

**WVE-006 therapeutic goal is to restore dynamic AAT physiology;
augmentation therapy goal is to maximize AAT levels as dynamic response is not enabled**

WVE-006 achieved key treatment goals of restoring MZ phenotype

Total AAT levels exceeded 11 μM , production of wild-type M-AAT of greater than 50%, restored physiological AAT production

Plasma AAT of ~13 μM

- Protein levels associated with lower risk of AATD liver and lung diseases

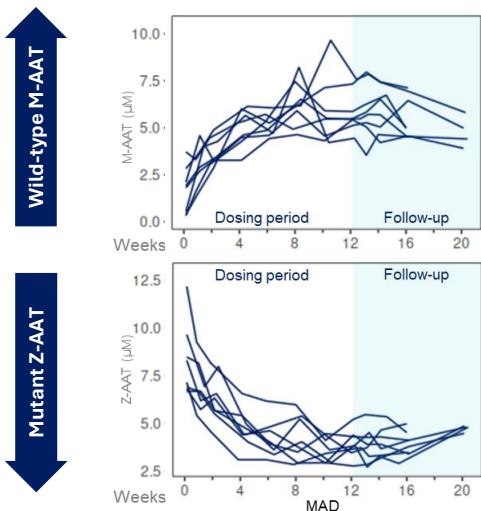
400 mg single dose

12.8 μM total AAT

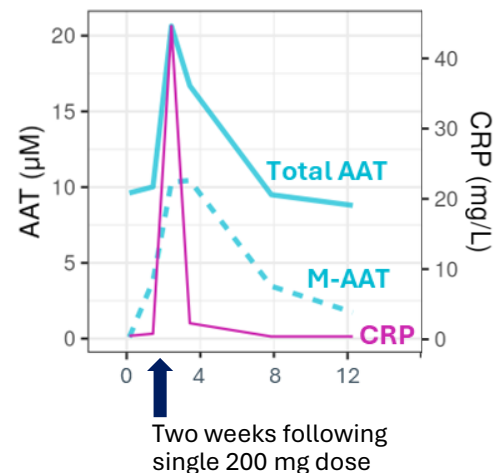
200 mg bi-weekly

11.9 μM total AAT

Wild-type M-AAT protein of 64% of total, reduction in Z-AAT



AAT reached >20 μM during an acute phase response



400 mg monthly and 600 mg single dose data expected in May 2026

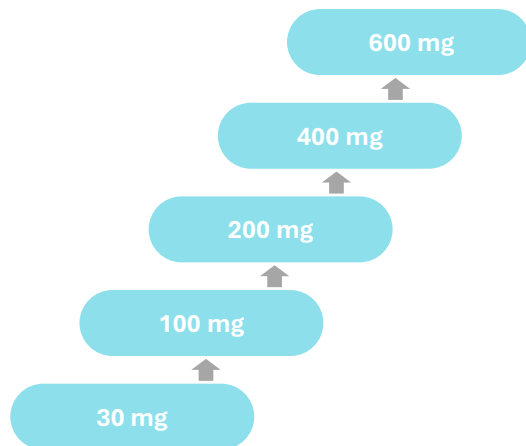
RestorAATion-2 clinical trial ongoing

RESTOR**AAT**ION

RestorAATion-1: Healthy Volunteers

RestorAATion-2: AATD Patients

SAD → MAD Multi-dosing complete



SAD Cohort 3
600 mg



MAD Cohort 3
600 mg; Q4W

SAD Cohort 2
400 mg



MAD Cohort 2
400 mg; Q4W

SAD Cohort 1
200 mg



MAD Cohort 1
200 mg Q2W

Study key objectives

Safety and tolerability

Pharmacokinetics

Serum M-AAT levels

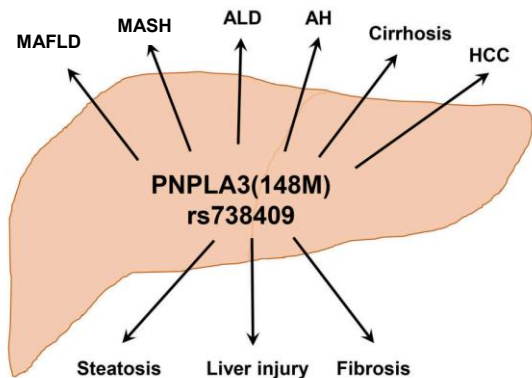
WVE-008
RNA editing (AlMer)

PNPLA3 I148M liver disease

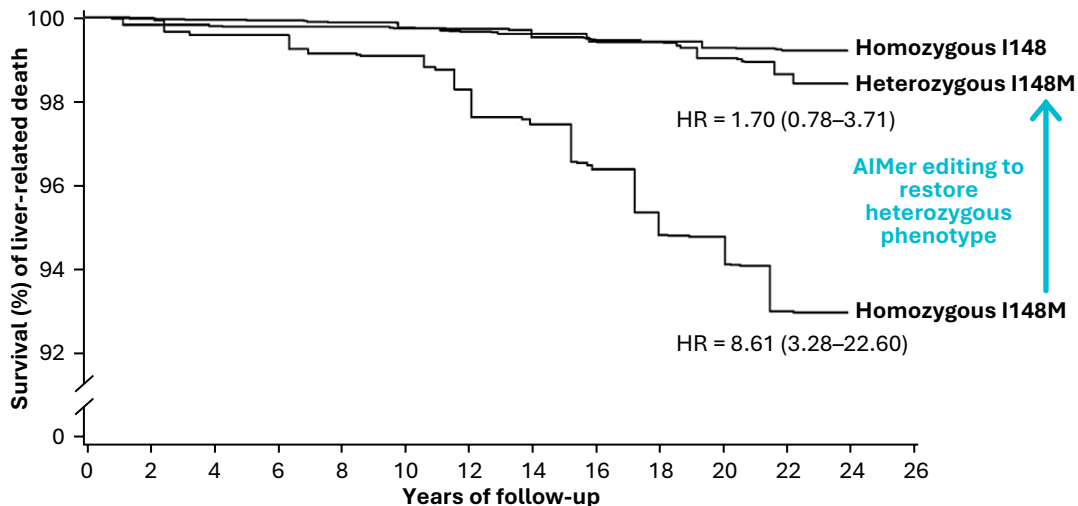
WVE-008 for PNPLA3 I148M liver disease

GalNAc-RNA editing approach uniquely aims to restore PNPLA3 function to fully address disease

Homozygous PNPLA3 I148M carriers have significantly higher risk of multiple liver diseases



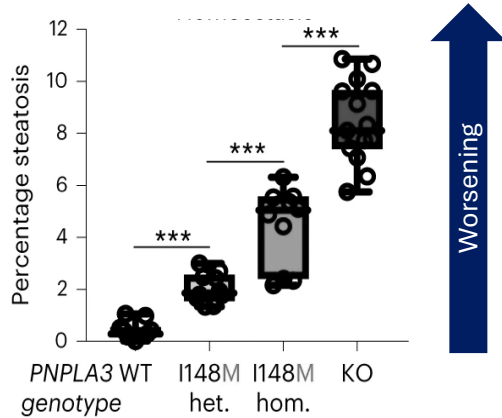
Heterozygous carriers have 80% lower risk of liver-related death as compared to homozygous carriers



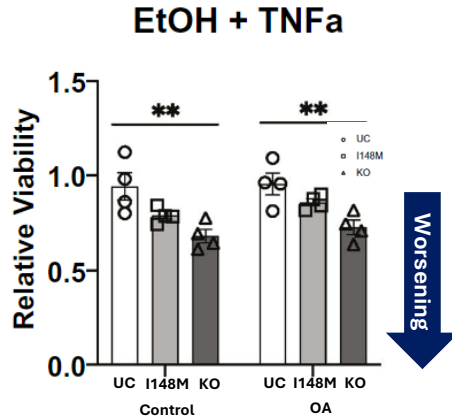
Over nine million homozygous PNPLA3 I148M patients with liver disease in US and Europe

Silencing of PNPLA3 in normal liver may worsen basal physiological functions

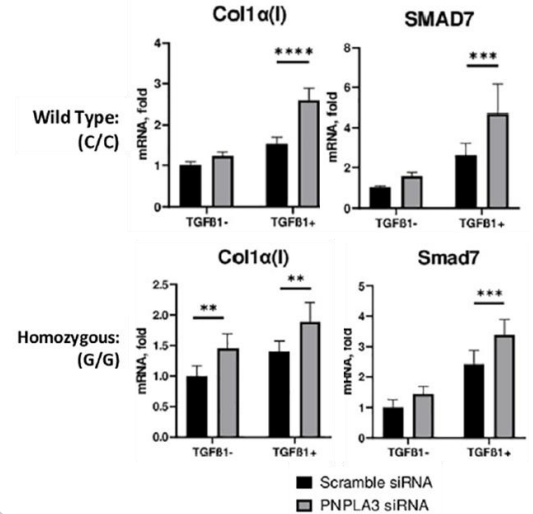
Silencing PNPLA3 worsens steatosis in iPSC-derived human liver organoids²



Silencing PNPLA3 increases inflammation-induced liver cell death in human primary hepatocytes³



PNPLA3 siRNA exacerbates the fibrotic response in hepatic stellate cells¹

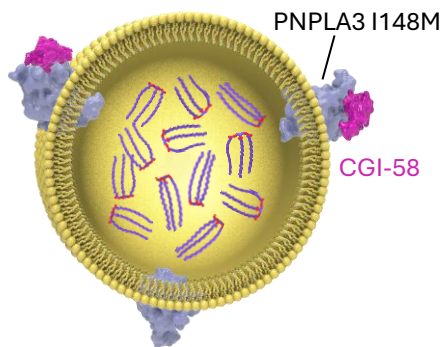


Functional PNPLA3 is imperative for liver health beyond improvements in steatosis

RNA editing is expected to restore PNPLA3 function to treat across the stages of liver diseases

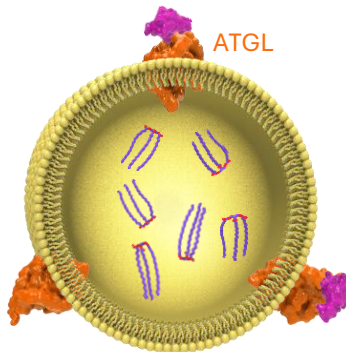
✓ RNA editing approach

PNPLA3 I148M aggravates steatosis and fibrosis through gain-of-function



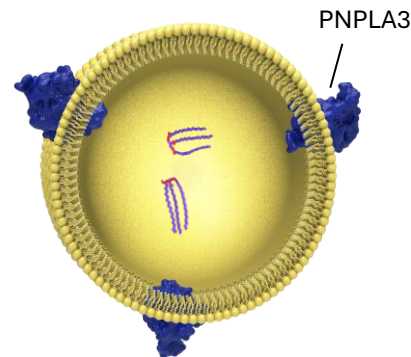
- PNPLA3 I148M accumulates on LDs, sequesters CGI-58, inhibits ATGL's lipase activity and lipid mobilization from ER
- Suppresses retinol metabolism in liver and worsens inflammation and fibrosis
- Promotes liver fat accumulation and fibrosis through activation of stellate cells

Silencing PNPLA3 may only partially address disease



- Creates PNPLA3 loss of function
- ATGL partial rescue for loss PNPLA3
- Silencing will not restore retinol metabolism
- **Fibrosis, ballooning, and inflammation persist**

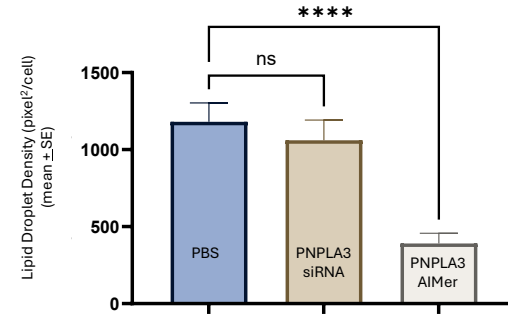
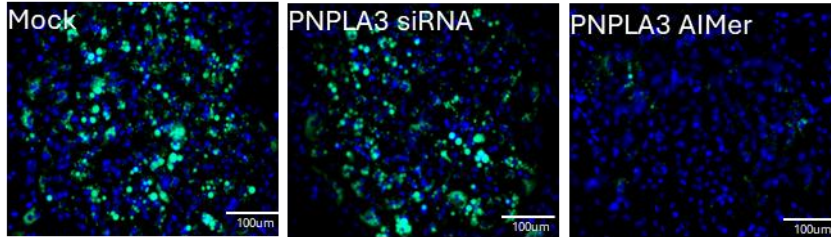
PNPLA3 correction expected to restore function, counter liver disease



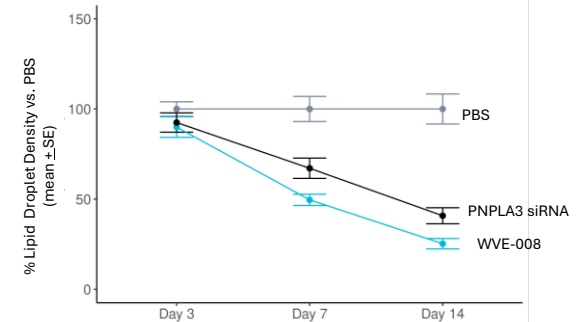
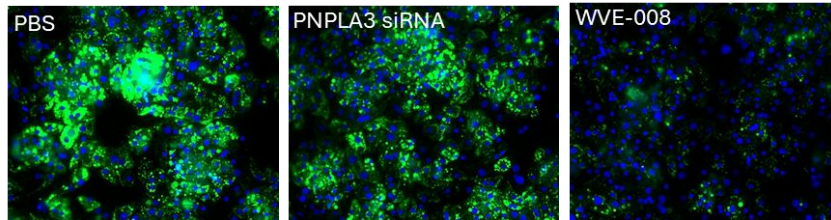
- Restores full PNPLA3 activity
- **Restores lipid mobilization, reverses steatosis, fibrosis, ballooning, and inflammation**

AIMers achieve efficient editing of PNPLA3, leading to reduction of liver fat

Significant decrease in liver fat with PNPLA3 editing in human HEPATOPAC® model with homozygous I148M

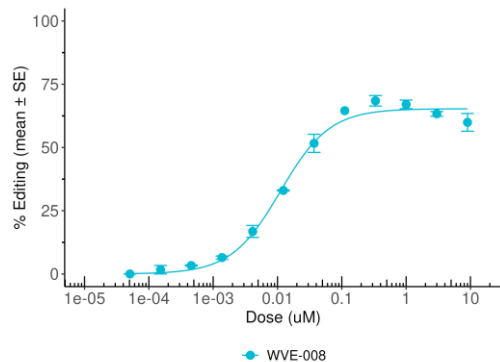


Decrease in liver fat with WVE-008 in monolayer model

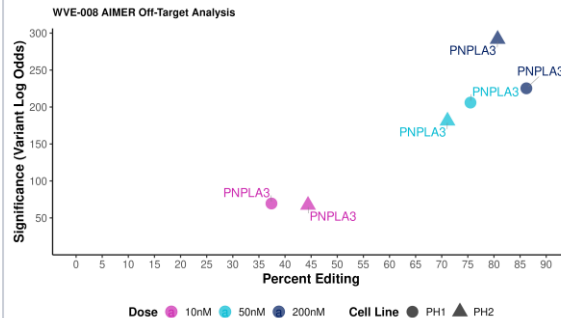


Preclinical data support WVE-008 as potential first-in-class, disease modifying therapy, for treatment of PNPLA3 I148M liver disease

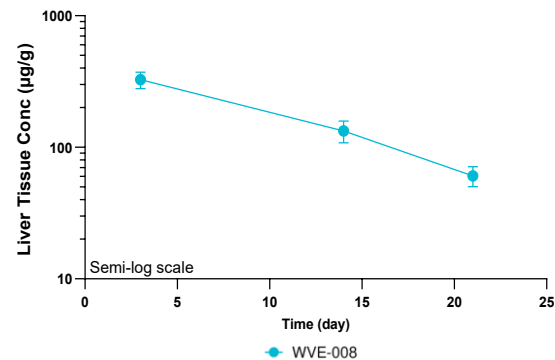
Potent editing with WVE-008



Highly specific editing with WVE-008



Tissue exposure supports excellent delivery

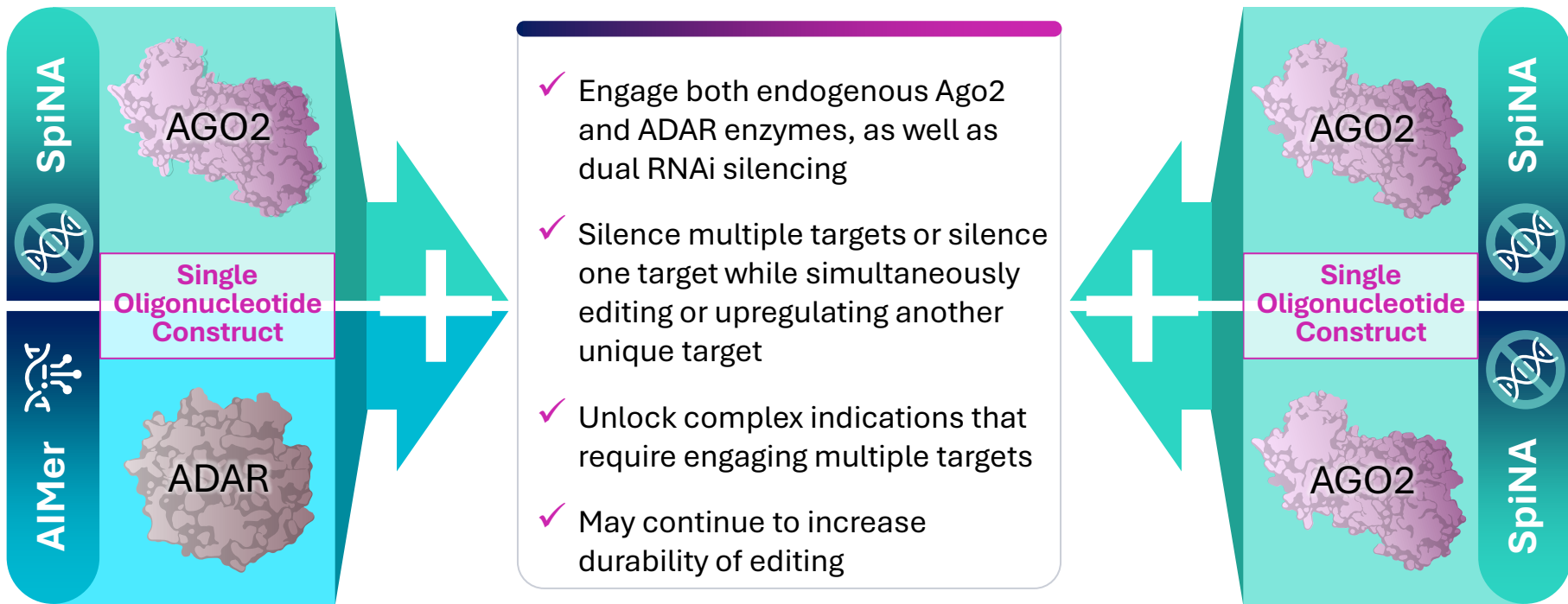


Expect to file Clinical Trial Application (CTA) for WVE-008 in 2026

Bifunctional modalities

Single oligonucleotide constructs

Reimagining RNA medicines: Bifunctional modalities



Other clinical programs

Duchenne muscular dystrophy

Advancing WVE-N531 in exon 53 amenable DMD

WVE-N531: exon skipping oligonucleotide designed to induce production of endogenous, functional dystrophin protein

- High unmet need for therapies delivering **more consistent dystrophin expression**, as few patients today achieve dystrophin >5% of normal
- **Opportunity to extend dosing intervals** beyond weekly standard of care to alleviate burden for patients and caregivers
- **Need to reach stem cells and distribute broadly to muscle tissues** to potentially enable muscle regeneration and impact respiratory and cardiac function
- WVE-N531 has Rare Pediatric Disease Designation and Orphan Drug Designation from FDA

DMD impacts ~1 / 5,000 newborn boys annually; ~20,000 new cases annually worldwide



FORWARD-53 48-week clinical trial results: WVE-N531's potential best-in-class profile for boys amenable to exon 53 skipping

- ✓ Statistically significant and clinically meaningful improvement (3.8s) in Time-to-Rise vs. natural history; functional benefits on other measures including NSAA
- ✓ Statistically significant reductions in muscle fibrosis and CK; driven by decreases in inflammation and necrosis; transition from regenerative to mature muscle
- ✓ Consistent dystrophin expression averaged 7.8% between 24 and 48 weeks, with 88% of boys above 5% dystrophin; delivery to both myofibers and muscle stem cells
- ✓ WVE-N531 remains generally safe and well-tolerated with no Serious Adverse Events

NDA filing for accelerated approval with monthly dosing planned for 2026

Reimagining RNA medicines

Poised for significant and sustained growth driven by RNAi and RNA editing



RNAi
SpiNA

WVE-007
Obesity

Other hepatic targets

Extra-hepatic targets

**Bifunctional single oligonucleotide
constructs**

Other hepatic targets

Extra-hepatic targets



**RNA
Editing**
AIMers

WVE-006
AATD
WVE-008
PNPLA3 I148M liver
disease

Anticipated upcoming milestones

WVE-007

(*INHBE*)

Obesity

- Deliver additional data from INLIGHT, including data from the 600 mg SAD cohort in 2026
- Initiate Phase 2a multidose portion of INLIGHT in individuals living with obesity with higher BMI with and without type 2 diabetes in 2Q 2026
- Combination and maintenance studies of WVE-007 expected to initiate in 2026

INLIGHT

WVE-006

(*SERPINA1*)

AATD

- Deliver data from 400 mg MAD cohort and 600 mg SAD cohort in May 2026
- Deliver multidose data from 600 mg cohort in 2H 2026
- Regulatory feedback on potential accelerated approval pathway expected mid-2026

RESTORAACTION

WVE-008

PNPLA3 I148M liver disease

- File CTA for WVE-008 in 2026

WVE-N531

(*exon 53*)

DMD

- Submit NDA to support accelerated approval of WVE-N531 with monthly dosing in 2026



For questions contact:
investorrelations@wavelifesci.com